



09/756,096, 09/158,863

# SEQUENCE LISTING

<110> Mitchell, Lloyd G.  
Garcia-Blanco, Mariano A.  
Puttaraju, Madaiah  
Mansfield, Gary S.

<120> METHODS AND COMPOSITIONS FOR USE IN  
SPLICEOSOME MEDIATED RNA TRANS-SPLICING

<130> A31304-B-A-B 072874.0135

<140> 09/756,096

<141> 2001-01-08

<150> 09/158,863

<151> 1998-09-23

<150> 09/133,717

<151> 1998-08-13

<150> 09/087,233

<151> 1998-05-28

<150> 08/766,354

<151> 1996-12-13

<150> 60/008,317

<151> 1995-12-15

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120

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132

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 24

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<400> 12  
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<400> 18

gttctgtcct tgtctc  
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<210> 19

<211> 29

<212> DNA

<213> Corynebacterium diptheriae

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ggcgctgcag ggcgctgatg atgttggtg  
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<210> 20

<211> 36

<212> DNA

<213> Corynebacterium diptheriae

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36

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<400> 23  
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<210> 24  
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<210> 27  
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<400> 27  
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 19

<210> 28  
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<210> 29  
 <211> 36  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Oligonucleotide primer complimentary to the  
 Escherichia coli lacZ gene

<400> 29  
 ctgaggatcc tcttacctgt aaacgcccac actgac  
 36

<210> 30  
 <211> 38  
 <212> DNA  
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<220>

<223> Oligonucleotide primer complimentary to the  
Escherichia coli lacZ gene

<400> 30

gcatggtaac cctgcagggc ggcttcgtct gggactgg  
38

<210> 31

<211> 38

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer complimentary to the  
Escherichia coli lacZ gene

<400> 31

ctgaaagctt gttaacttat tatttttgac accagacc  
38

<210> 32

<211> 47

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer complimentary to the  
Escherichia coli lacZ gene

<400> 32

gcatggtaac cctgcagggc ggcttcgtct aataatggga ctgggtg  
47

<210> 33

<211> 37

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer complimentary to the beta  
HCG6 gene (accession #X00266)

<400> 33

gcatggatcc tccggagggc ccctgggcac ctccac  
37

<210> 34



<211> 38

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer complimentary to the beta  
HCG6 gene (accession #X00266)

<400> 34

ctgactgcag ggtaaccgga caaggacact gcttcacc  
38

<210> 35

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer complimentary to the beta  
HCG6 gene (accession #X00266)

<400> 35

gcatggtaac cctgcagggg ctgctgctgt tgctg  
35

<210> 36

<211> 37

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer complimentary to the beta  
HCG6 gene (accession #X00266)

<400> 36

ctgaaagctt gttaaccagc tcacccatggt ggggcag  
37

<210> 37

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer complimentary to the  
Escherichia coli lacZ gene

<400> 37

ggcttttcgct acctggagag ac  
22

<210> 38  
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<220>  
<223> Oligonucleotide primer complimentary to the  
Escherichia coli lacZ gene

<400> 38  
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21

<210> 39  
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<212> DNA  
<213> Artificial Sequence

<220>  
<223> Oligonucleotide primer complimentary to the  
Escherichia coli lacZ gene

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20

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<400> 40  
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45

<210> 41  
<211> 35  
<212> DNA  
<213> Homo sapiens

<400> 41  
acctctgcag gtgaccctgc aggaaaaaaa agaag  
35

<210> 42

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<211> 30
<212> DNA
<213> Homo sapiens
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<400> 42
acctctgcag acttcacttc taatgatgat
30
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<210> 43
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<212> DNA
<213> Homo sapien
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51
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<210> 44
<211> 32
<212> DNA
<213> Homo sapien
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<400> 44
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32
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<210> 45
<211> 35
<212> DNA
<213> Homo sapien
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<400> 45  
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35

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<210> 46
<211> 35
<212> DNA
<213> Homo sapien
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<210> 47
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<213> Homo sapien
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<400> 47  
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32

<210> 48  
<211> 21  
<212> DNA  
<213> Homo sapien

<400> 48  
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21

<210> 49  
<211> 21  
<212> DNA  
<213> Homo sapien

<400> 49  
cgcctaatga tgatgatgat g  
21

<210> 50  
<211> 21  
<212> DNA  
<213> Homo sapien

<400> 50  
cttcttggtgta ctctgtcct g  
21

<210> 51  
<211> 32  
<212> DNA  
<213> Homo sapien

<400> 51  
gacctctcga gggatttggg gaattatttg ag  
32

<210> 52  
<211> 21  
<212> DNA  
<213> Homo sapien

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aactagaagg cacagtcgag g



according to specification

<400> 55

gcuagccugg gacaaggaca cugcuucacc cgguuaguag accacagccc ugagccnnnn  
60  
nnnnnnnnnn aucguuaacu aaauaacuac uaacuggggug aacuucugua uuauucucga  
120  
gcugcag  
127

<210> 56

<211> 127

<212> RNA

<213> Artificial Sequence

<220>

<223> PTM intramolecular base paired stem

<221> misc\_feature

<222> (57)...(70)

<223> Loop comprising a combination of 14 nucleotides  
according to specification

<400> 56

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nnnnnnnnnn aucguuaacu aaauaacuac uaacuggggug aaguucuguc cuugucucga  
120  
gcugcag  
127

<210> 57

<211> 132

<212> DNA

<213> Artificial Sequence

<220>

<223> trans-spliced product containing Human chorionic  
gonadotropin gene 6 sequences and Corynebacterium  
diphtheriae diphtheria toxin A sequences

<400> 57

caggggacgc accaaggatg gagatgttcc agggcgctga tgatgttggtt gattcttctt  
60  
aaatcttttg tgatggaaaa cttttcttcg taccacggga ctaaacttgg ttatgtagat  
120  
tccattcaaa aa  
132

<210> 58  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Artificial Sequence derived from Escherichia coli  
lacZ gene

<400> 58  
gaattcggta ccatgggg  
18

<210> 59  
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<220>  
<223> Artificial Sequence derived from Escherichia coli  
lacZ gene

<400> 59  
cgtttacagg taagaggatc ctccggaggg ccc  
33

<210> 60  
<211> 30  
<212> DNA  
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<220>  
<223> Artificial Sequence derived from Escherichia coli  
lacZ gene

<400> 60  
tggtgtcaaa aataataagt taacaagctt  
30

<210> 61  
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<212> DNA  
<213> Artificial Sequence

<220>  
<223> trans-spliced product containing Escherichia coli  
lacZ gene sequences and Human chorionic

gonadotropin gene 6 exon 2 sequences

<400> 61

cagcagcccc tgtaaacggg gatac  
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<210> 62

<211> 286

<212> DNA

<213> Artificial Sequence

<220>

<223> trans-spliced product containing Escherichia coli  
lacZ gene sequences

<400> 62

ggctttcgct acctggagag acgcgcccgc tgatcctttg cgaatacgcc cacgcgatgg  
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gtaacagtct tggcgggtttc gctaaatact ggcaggcggt tcgtcagtat ccccgtttac  
120  
agggcggtt cgtctaataa tgggactggg tggatcagtc gctgattaaa tatgatgaaa  
180  
acgggcaacc cgtggtcggc ttacggcggt gatcttggcg atacgccgaa cgatcgccag  
240  
ttctgtatga acggtctggt ctttgccgac cgcacgccc atccag  
286

<210> 63

<211> 196

<212> DNA

<213> Artificial Sequence

<220>

<223> trans-spliced product containing Escherichia coli  
lacZ gene sequences

<400> 63

ggctttcgct acctggagag acgcgcccgc tgatcctttg cgaatacgcc cacgcgatgg  
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gtaacagtct tggcgggtttc gctaaatact ggcaggcggt tcgtcagtat ccccgtttac  
120  
aggggctgct gctggttctg ctgctgagca tgggcgggac atgggcatcc aaggagccac  
180  
ttcggccacg gtgcg  
196

<210> 64

<211> 420



<212> DNA

<213> Artificial Sequence

 $\langle 220 \rangle$ 

<223> trans-spliced product comprising cystic fibrosis  
transmembrane regulator-derived sequences and His  
tag sequence

<400> 64

gctagcgttt aaacgggccc acccatcatt attaggtcat tatccgcgga acattattat  
60

aacgttgctc gagtactaac tggaacctct tctttttttt cctgcagact tcacttctaa  
120

tgatgattat gggagaactg gagccttcag agggtaaaat taagcacagt ggaagaattt  
180

cattctgttc tcagttttcc tggattatgc ctggcaccat taaagaaaat atcatctttg  
240

cgggccgcca ctgtgctgga tatctgcaga attccaccac actggactag tggatccgag  
300

ctcggtacca aggttaagtt taaaccgctg atcagcctcg actgtgcctt ctagttgcc  
360

gccatctggt gtttgccctt ccccggtgcc ttcttgacc ctggaagggt ccaactccac  
420

<210> 65

$\langle 211 \rangle$  20

<212> DNA

<213> Artificial Sequence

 $\langle 220 \rangle$ 

<223> Splice junction sequence

<400> 65

atgttccagg gcgtgatgat

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<210> 66

<211> 7

&lt;212&gt; PRT

<213> Artificial Sequence

 $\langle 220 \rangle$ 

<223> C terminal residues from glutathione -S-transferase

<400> 66

Asp Tyr Lys Asp Asp Asp Lys

1

5

<210> 67  
<211> 15  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Artificial sequence comprising sequences derived  
from Escherichia coli lacZ gene

<400> 67  
ggagttgatc ccgtc  
15

<210> 68  
<211> 37  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Artificial sequence comprising sequences derived  
from Escherichia coli lacZ gene

<400> 68  
gcagtgtcct tgtgcgggta ccctgcaggg cggttc  
37

<210> 69  
<211> 120  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Binding domain of PTM

<400> 69  
gattcacttg ctccaattat catcctaagc agaagtgtat attcttattt gtaaagattc  
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tattaactca ttgattcaa aatatttaaa atacttcttg ttccatactc tgctatgcac  
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<210> 70  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Spacer sequence of PTM

<400> 70

aacattatta taacggttgct cgaa  
24

<210> 71

<211> 47

<212> DNA

<213> Artificial Sequence

<220>

<223> Branch point, pyrimidine tract and acceptor splice  
site of PTM

<400> 71

tactaactgg tacctcttct tttttttttg atatacctgca gggcggc  
47

<210> 72

<211> 70

<212> DNA

<213> Artificial Sequence

<220>

<223> Donor site and spacer sequence of PTM

<400> 72

tgaacggtaa gtgttatcac cgatatgtgt ctaacctgat tcgggccttc gatacgctaa  
60  
gatccaccgg  
70

<210> 73

<211> 260

<212> DNA

<213> Artificial Sequence

<220>

<223> Binding domain of spacer sequence

<400> 73

tcaaaaagtt ttcacataat ttcttacctc ttcttgaatt catgctttga tgacgcttct  
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120



<220>

<223> Oligonucleotide primer

<400> 77

ctaggatccc gttcttttgt tcttcactat taa  
33

<210> 78

<211> 33

<212> DNA

<213> Artificial Sequence

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<223> Oligonucleotide primer

<400> 78

ctagggttac cgaagtaaaa ccatacttat tag  
33

<210> 79

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer

<400> 79

gcatgggttac cctgcagggg ctgctgctgt tgctg  
35

<210> 80

<211> 37

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer

<400> 80

ctgaaagctt gttaaccagc tcaccatggt ggggcag  
37

<210> 81

<211> 23

<212> DNA

<213> Artificial Sequence

<220>  
<223> Binding domain of PTM molecule

<400> 81  
accatcatt attaggtcat tat  
23

<210> 82  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Oligonucleotide primer

<400> 82  
gatcaaattct gtcgatacctt cc  
22

<210> 83  
<211> 21  
<212> DNA  
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<220>  
<223> Oligonucleotide primer

<400> 83  
ctgatccacc cagtccatt a  
21

<210> 84  
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<400> 84  
gactgatcca cccagtccca ga  
22

<210> 85  
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<223> Random sequence inserted to replace 3' splice site

<221> misc\_feature

<222> (7)...(30)

<223> spacer sequence, see SEQ ID NO 70

<400> 85

ccgcggnnnn nnnnnnnnnn nnnnnnnnnn gggttccggt accggcggct tc  
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<210> 86

<211> 71

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide

<400> 86

ttttatcccc gtttacaggg cggcttcgtc tgggactggg tggatcagtc gctgattaaa  
60  
tatgatgaaa a  
71

<210> 87

<211> 66

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide

<400> 87

tttggcgata cgccgaacga tcgccagttc tgtatgaacg gtctgggtctt tgccgaccgc  
60  
acgccg  
66

<210> 88

<211> 192

<212> DNA

<213> Artificial Sequence

<220>

<223> PTM sequences

<400> 88

acgagcttgc tcatgatgat catgggagag ttagaaccaa gtgaaggcaa gatcaaaca  
60  
tccggccgca tcagcttttg cagccaattc agttggatca tgcccggtag catcaaggag  
120  
aacataatct tcggcgtagc ttacgacgag taccgctatc gctcggtagc taaggcctgt  
180  
cagttggagg ag  
192

<210> 89  
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<212> DNA  
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<220>  
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<400> 89  
gagcaggcaa gacgagcttg ctcac  
25

<210> 90  
<211> 28  
<212> DNA  
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<220>  
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<400> 90  
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28

<210> 91  
<211> 30  
<212> DNA  
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<220>  
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<400> 91  
gtcagttgga ggaggacatc tccaagtttg  
30

<210> 92  
<211> 192  
<212> DNA



<213> Artificial Sequence

<400> 92

acgagcttgc tcatgatgat catgggagcgag ttagaaccaa gtgaaggcaa gatcaaacat  
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tccggcgcga tcagcttttg cagccaattc agttggatca tgcccggtag catcaaggag  
120  
aacataatct tcggcgtcag ttacgacgag taccgctatc gctcgggtgat taaggcctgt  
180  
cagttggagg ag  
192

<210> 93

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> PTM sequences

<400> 93

aaatatcatt ggtgttttctt atgatga  
27

<210> 94

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide

<400> 94

ccaactagaa gaggacatct ccaagtttgc  
30

<210> 95

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide

<400> 95

atgatcatgg gcgagttaga accaagttag  
30

<210> 96  
 <211> 27  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Oligonucleotide

<400> 96  
 aaaatatcat ctttggtggt tcctatg  
 27

<210> 97  
 <211> 27  
 <212> DNA  
 <213> Artificial Sequence

<220>  
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<400> 97  
 ccaactagaa gaggacatct ccaagtt  
 27

<210> 98  
 <211> 21  
 <212> DNA  
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<220>  
 <223> 5' splice site

<400> 98  
 cgtttacagg taagtggatc c  
 21

<210> 99  
 <211> 27  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> 3' splice site

<400> 99  
 ctgcagggcg gcttcgtcta ataatgg  
 27

<210> 100  
 <211> 47  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Sequence from trans-splicing domain

<400> 100  
 tactaactgg tacctcttct tttttttttg atatcctgca gggcggc  
 47

<210> 101  
 <211> 1584  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> CFTR PTM

<400> 101  
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 agaccaatth tgaggaaagg atacagacag cgcctggaat tgtcagacat ataccaaath  
 120  
 ccttctgttg attctgctga caatctatct gaaaaattgg aaagagaath ggatagagag  
 180  
 ctggcttcaa agaaaaatcc taaactcatt aatgcccttc ggcatgttt tttctggaga  
 240  
 tttatgttct atggaatctt tttatattha ggggaagtca ccaaagcagt acagcctctc  
 300  
 ttactgggaa gaatcatagc ttctatgac ccggataaca aggaggaacg ctctatcgcg  
 360  
 atttatctag gcataggctt atgccttctc tttattgtga ggacactgct cctacacca  
 420  
 gccatttttg gccttcatca cattggaath cagatgagaa tagctatgtt tagtttgatt  
 480  
 tataagaaga ctttaaagct gtcaagccgt gttctagata aaataagtat tggacaactt  
 540  
 gttagtctcc tttccaacaa cctgaacaaa tttgatgaag gacttgcatt ggcacatttc  
 600  
 gtgtggatcg ctcttttgca agtggcactc ctcatggggc taatctggga gttgttacag  
 660  
 gcgtctgcct tctgtggact tggtttcctg atagtccttg ccctttttca ggctgggcta  
 720  
 gggagaathg tgatgaagta cagagatcag agagctggga agatcagtha aagacttgth  
 780  
 attacctcag aathgatcga gaacatccaa tctgtthaagg catactgctg ggaagaagca

840  
atggaaaaaa tgattgaaaa ctttaagacaa acagaactga aactgactcg gaaggcagcc  
900  
tatgtgagat acttcaatag ctcagccttc ttcttctcag ggttctttgt ggtgttttta  
960  
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1020  
tcattctgca ttgttctgcg catggcggtc actcggcaat ttccctgggc tgtacaaaca  
1080  
tggtatgact ctcttgagc aataaacaaa atacaggatt tcttacaaaa gcaagaatat  
1140  
aagacattgg aatataactt aacgactaca gaagtagtga tggagaatgt aacagccttc  
1200  
tgggaggagg gatttgggga attatttgag aaagcaaaac aaaacaataa caatagaaaa  
1260  
acttctaata gtgatgacag cctcttcttc agtaatttct cacttcttgg tactcctgtc  
1320  
ctgaaagata ttaatttcaa gatagaaaga ggacagttgt tggcggttgc tggatccact  
1380  
ggagcaggca agacgagctt gctcatgatg atcatgggag agttagaacc aagtgaaggc  
1440  
aagatcaaac attccggccg catcagcttt tgcagccaat tcagttggat catgcccggg  
1500  
accatcaagg agaacataat cttcggcgtc agttacgacg agtaccgcta tcgctcggtg  
1560  
attaaggcct gtcagttgga ggag  
1584

<210> 102  
<211> 323  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> trans-splicing domain of CFTR PTM

<400> 102  
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ggtcaaaaag ttttcacata atttcttacc tcttcttgaa ttcattgcttt gatgacgctt  
120  
ctgtatctat attcatcatt ggaaacacca atgatatttt ctttaattgg gcctggcata  
180  
atcctggaaa actgataaca caatgaaatt cttccactgt gcttaatttt accctctgaa  
240  
ttctccattt ctcccataat catcattaca actgaactct ggaaataaaa cccatcatta  
300  
ttaactcatt atcaaatcac gct

323

<210> 103  
 <211> 165  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> PTM binding domain

<400> 103  
 gctagcaata atgacgaagc cgccctcac gctcaggatt cacttgccctc caattatcat  
 60  
 cctaagcaga agtgtatatt cttatttgta aagattctat taactcattt gattcaaaat  
 120  
 atttaaaata cttcctgttt cacctactct gctatgcacc cgcg  
 165

<210> 104  
 <211> 225  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> trans-splicing domain of CFTR PTM

<400> 104  
 aataatgacg aagccgcccc tcacgctcag gattcacttg ccctccaatt atcatcctaa  
 60  
 gcagaagtgt atattcttat ttgtaaagat tctattaact catttgattc aaaatattta  
 120  
 aaataacttcc tgtttcacct actctgctat gcacccgcgg aacattatta taacgttgct  
 180  
 cgaataactaa ctggtacctc ttcttttttt tttgatatcc tgcag  
 225

<210> 105  
 <211> 3069  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> CFTR PTM sequence

<400> 105  
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 60  
 agtgggaagaa tttcattctg ttctcagttt tcctggatta tgctggcac cattaagaa

120  
 aatatcatct ttggtgtttc ctatgatgaa tatagataca gaagcgtcat caaagcatgc  
 180  
 caactagaag aggacatctc caagtgttgc gagaaagaca atatagttct tggagaaggt  
 240  
 ggaatcacac tgagtggagg tcaacgagca agaatttctt tagcaagagc agtatacaaa  
 300  
 gatgctgatt tgtatttatt agactctcct tttggatacc tagatgtttt aacagaaaaa  
 360  
 gaaatatttg aaagctgtgt ctgtaaactg atggctaaca aaactaggat tttggctact  
 420  
 tctaaaatgg aacatttaaa gaaagctgac aaaatattaa ttttgcataa aggtagcagc  
 480  
 tatttttatg ggacattttc agaactccaa aatctacagc cagactttag ctcaaaactc  
 540  
 atgggatgtg attctttcga ccaatttagt gcagaaagaa gaaattcaat cctaactgag  
 600  
 accttacacc gtttctcatt agaaggagat gctcctgtct cctggacaga aacaaaaaaa  
 660  
 caatctttta aacagactgg agagtgttgg gaaaaaagga agaattctat tctcaatcca  
 720  
 atcaactcta tacgaaaatt ttccattgtg caaaagactc ccttacaaat gaatggcatc  
 780  
 gaagaggatt ctgatgagcc tttagagaga aggctgtcct tagtaccaga ttctgagcag  
 840  
 ggagaggcga tactgcctcg catcagcgtg atcagcactg gccccacgct tcaggcacga  
 900  
 aggaggcagt ctgtcctgaa cctgatgaca cactcagtta accaagggtca gaacattcac  
 960  
 cgaaagacaa cagcatccac acgaaaagtg tcaactggccc ctcaggcaaa cttgactgaa  
 1020  
 ctggatatat attcaagaag gttatctcaa gaaactggct tggaaataag tgaagaaatt  
 1080  
 aacgaagaag acttaaagga gtgctttttt gatgatatgg agagcatacc agcagtgact  
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 1200  
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 1260  
 ggaaacactc ctcttcaaga caaagggaaat agtactcata gtagaaataa cagctatgca  
 1320  
 gtgattatca ccagcaccag ttcgtattat gtgttttaca tttacgtggg agtagccgac  
 1380  
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 1440  
 tcgaaaatth tacaccacaa aatgtttacat tctgtttctc aagcacctat gtcaaccctc  
 1500  
 aacacgttga aagcaggtgg gattcttaat agattctcca aagatatagc aattttggat

1560  
 gaccttctgc ctcttaccat atttgacttc atccagttgt tattaattgt gattggagct  
 1620  
 atagcagttg tcgcagtttt acaaccctac atctttgttg caacagtgcc agtgatagtg  
 1680  
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 1740  
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 1800  
 cttcgtgcct tcggacggca gccttacttt gaaactctgt tccacaaagc tctgaattta  
 1860  
 catactgcc actgggttctt gtacctgtca acactgcgct ggttccaaat gagaatagaa  
 1920  
 atgatttttg tcatcttctt cattgctggtt accttcattt ccattttaac aacaggagaa  
 1980  
 ggagaaggaa gagttggtat tatcctgact ttagccatga atatcatgag tacattgcag  
 2040  
 tgggctgtaa actccagcat agatgtggat agcttgatgc gatctgtgag ccgagtcctt  
 2100  
 aagttcattg acatgccaac agaaggtaaa cctaccaagt caaccaaacc atacaagaat  
 2160  
 ggccaactct cgaaagttat gattattgag aattcacacg tgaagaaaga tgacatctgg  
 2220  
 ccctcagggg gccaaatgac tgtcaaagat ctcacagcaa aatacacaga aggtggaaat  
 2280  
 gccatattag agaacatttc cttctcaata agtcctggcc agaggggtggg cctcttggga  
 2340  
 agaactggat caggaagag tactttgtta tcagcttttt tgagactact gaacactgaa  
 2400  
 ggagaaatcc agatcgatgg tgtgtcttgg gattcaataa ctttgcaaca gtggaggaaa  
 2460  
 gcctttggag tgataccaca gaaagtattt atttttctg gaacatttag aaaaaacttg  
 2520  
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 2580  
 agatctgtga tagaacagtt tcctgggaag cttgactttg tccttgtgga tgggggctgt  
 2640  
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 2880  
 gattccatcc agaaactgct gaacgagagg agcctcttcc ggcaagccat cagccctcc  
 2940  
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3000

3060

3069